



1/12

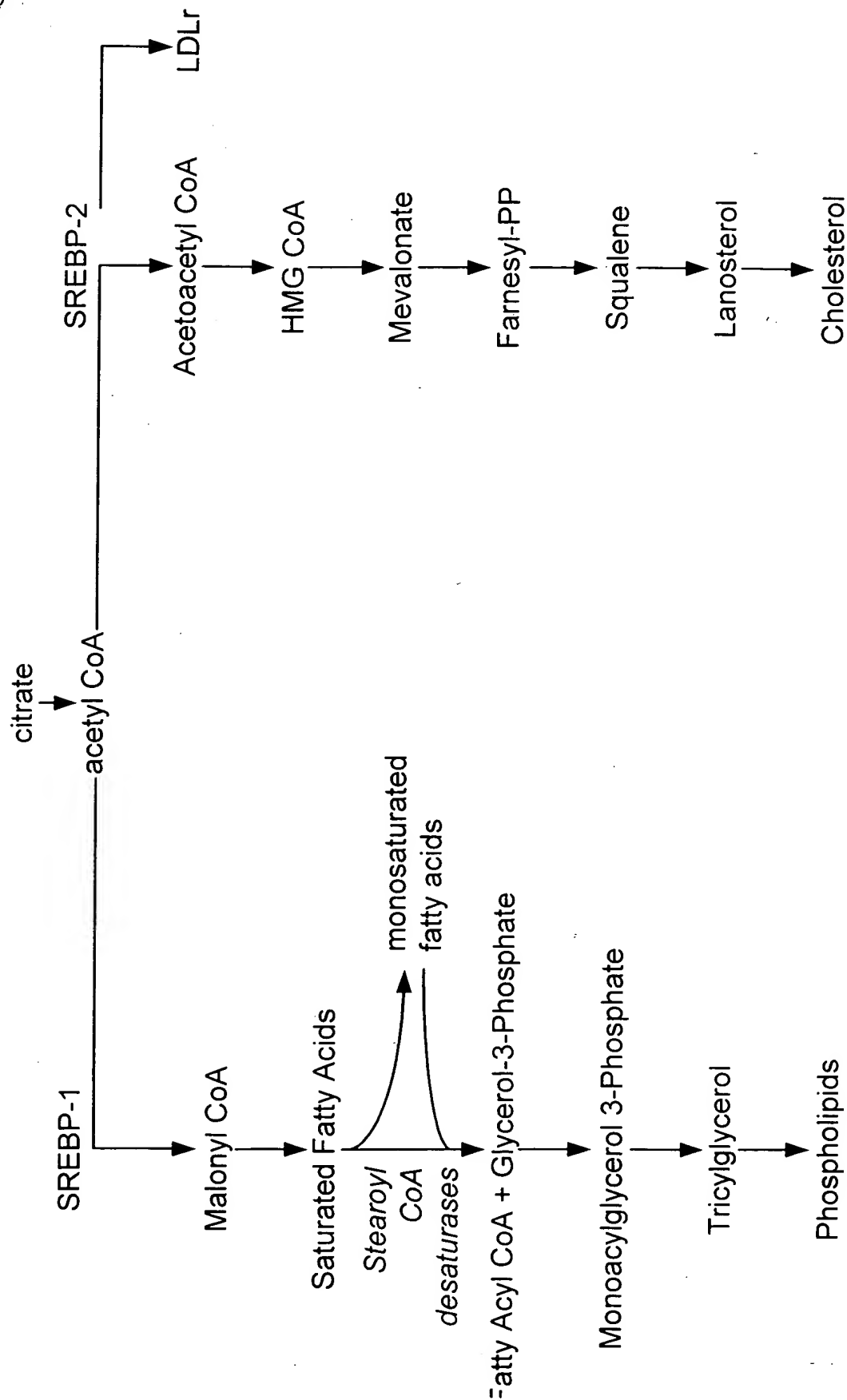


FIG. 1.

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Appl. No.: 09/848,990

Filed: May 3, 2001

TREATMENT OF HYPERTRIGLYCERIDEMIA AND OTHER
CONDITIONS USING LXR MODULATORS

Replacement Sheet

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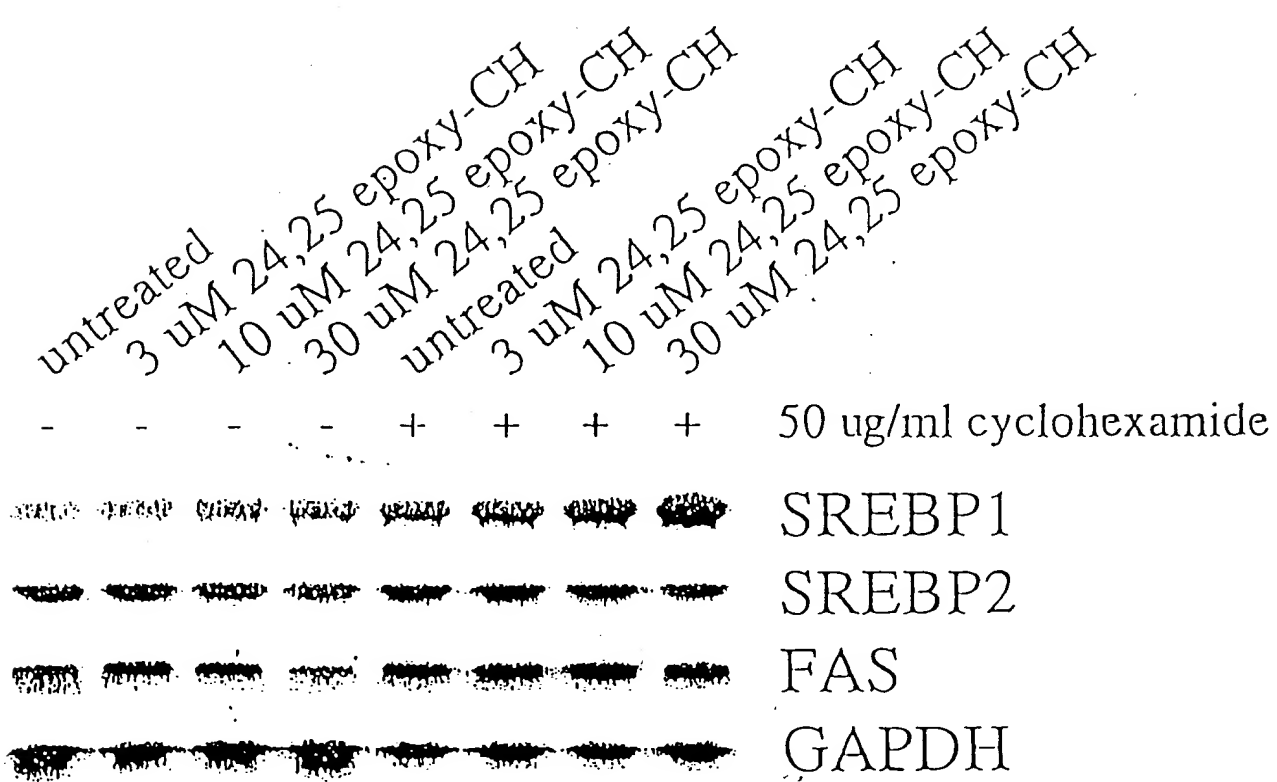


FIG. 2.

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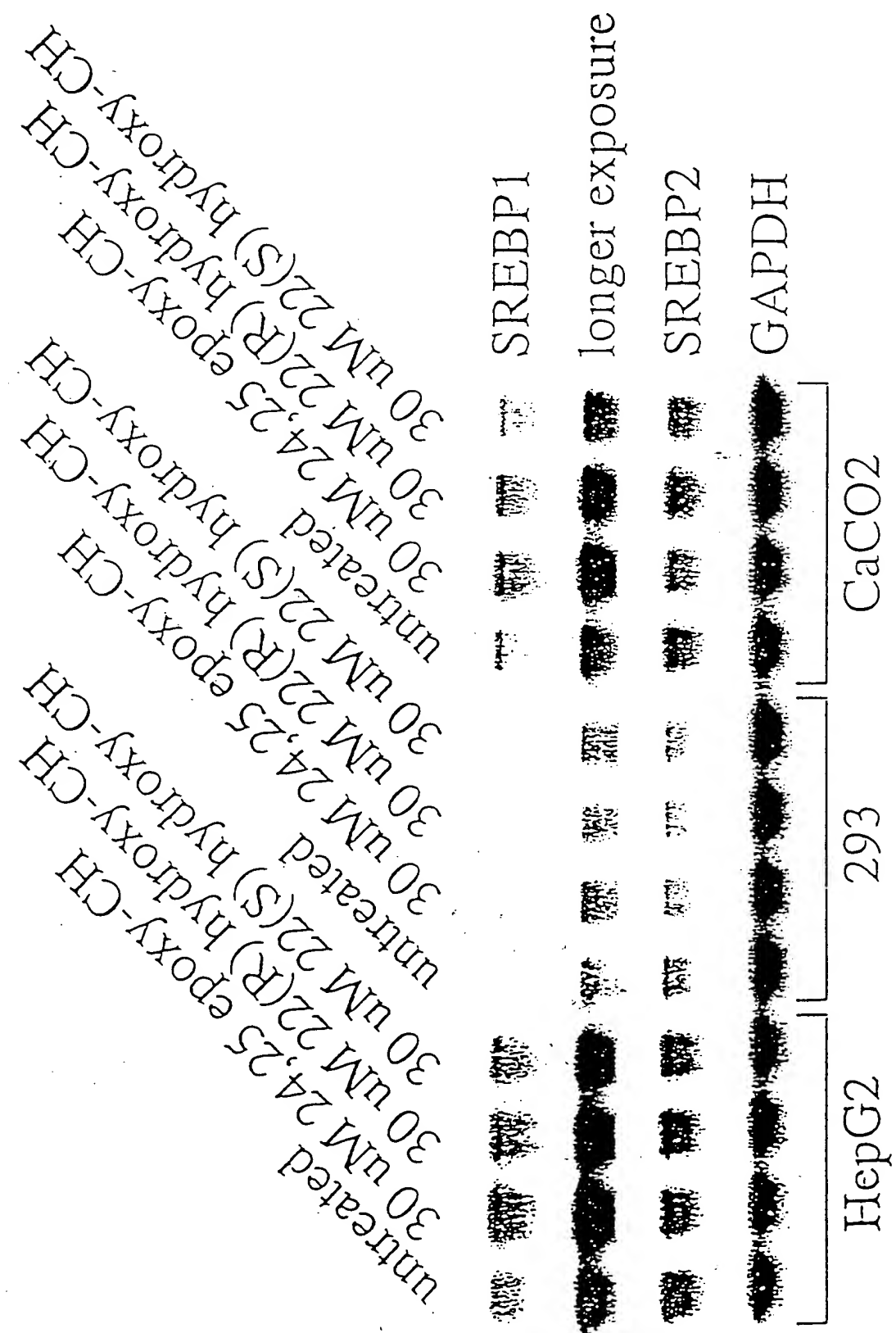


FIG. 3.

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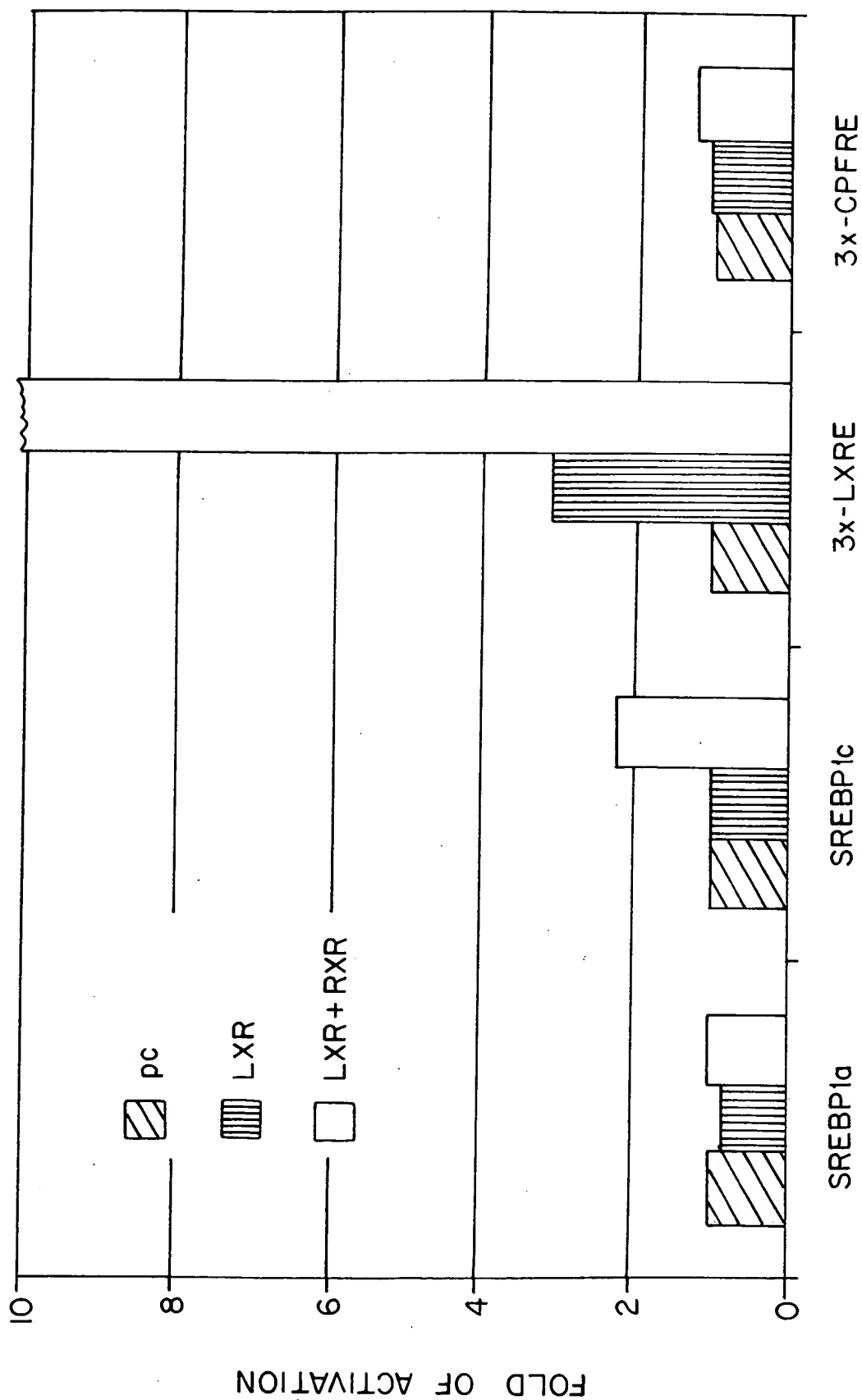
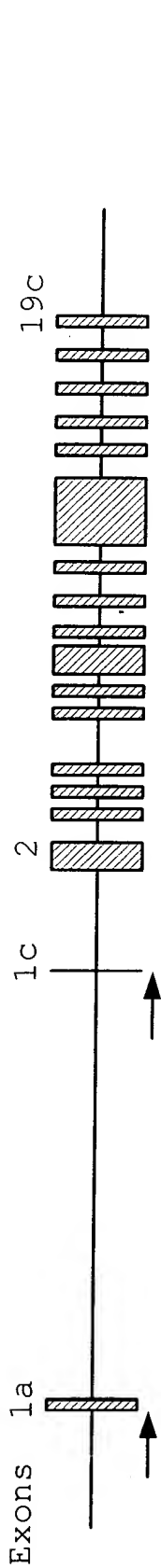


FIG. 4.

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in SREBP1c promoter

TGGGACCTGCCTGGCATGTGGAAGGGGCTTGGGGTGCAGCTGTGTCCCTGGCCCGCAGCACTGCCACACCCCTCGGT
TGACCTCACTTTTGGCTCTGGAGGGCAGAAACCTCTGAGCCAGCTCTCAGGGATACCTTTTTCCTCGATGGCCTT
GGAACTGCCCTCGCTATCTCTGAGCCCTGCCTGCCCTTCCCCCAAAGCCATCATAGACCCCTACCTCCTAT
ATGGAGCCTGGCCATGGAATGCTTGGGGTGAGAGGGACCCCATAGCGGCTCCGGGCATTGTTAGGGGGCTAGGTGG
TGCAGAGCCCCACCTGTGCGAGGGTTTACATGCCGCCACCCCTGCCCTCCTTGGGTAGCTGGGGTCAGAAAGGG
AGGCCTCACTCTCTGCATCTTAGTTTCTTACATCTAAAATGCTATTATATGCTGCTGAGGGTGGTTGGGG
CGTGCACTAGTGAGAGAGAGCCCTTCAGCCAGAGGACCCCATATGCTTAAACATCCAGACACCAAGGCCACTCA
CAGCAGCCGGCTTGTGTTCTGTCATCCCTTTTGGGTGAGGAACTGGGGACGCTGGACTCCAGATCTCCAGGAGGCC
CAGACTAGCGGGGAGGAGCCCTCAGCCCTGAGCTGAAACCTGAAATTTCTAGAATAAACCCAGGCCCTCCCTCCC
TGGGCAGCGGGGAGGGGTAGGGCCCAACGGCCTGGACGCCCCCAAGGGCGGCGAGATCGCGGAGCCATGGATTGCA

CGAAGgtatttttgaggcctccccac

FIG. 5.

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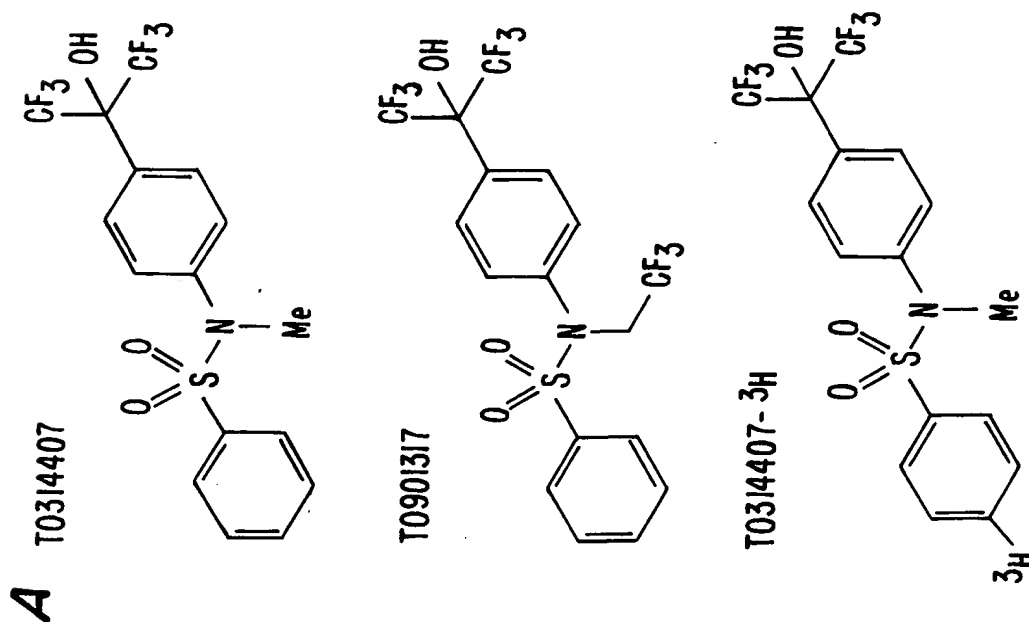
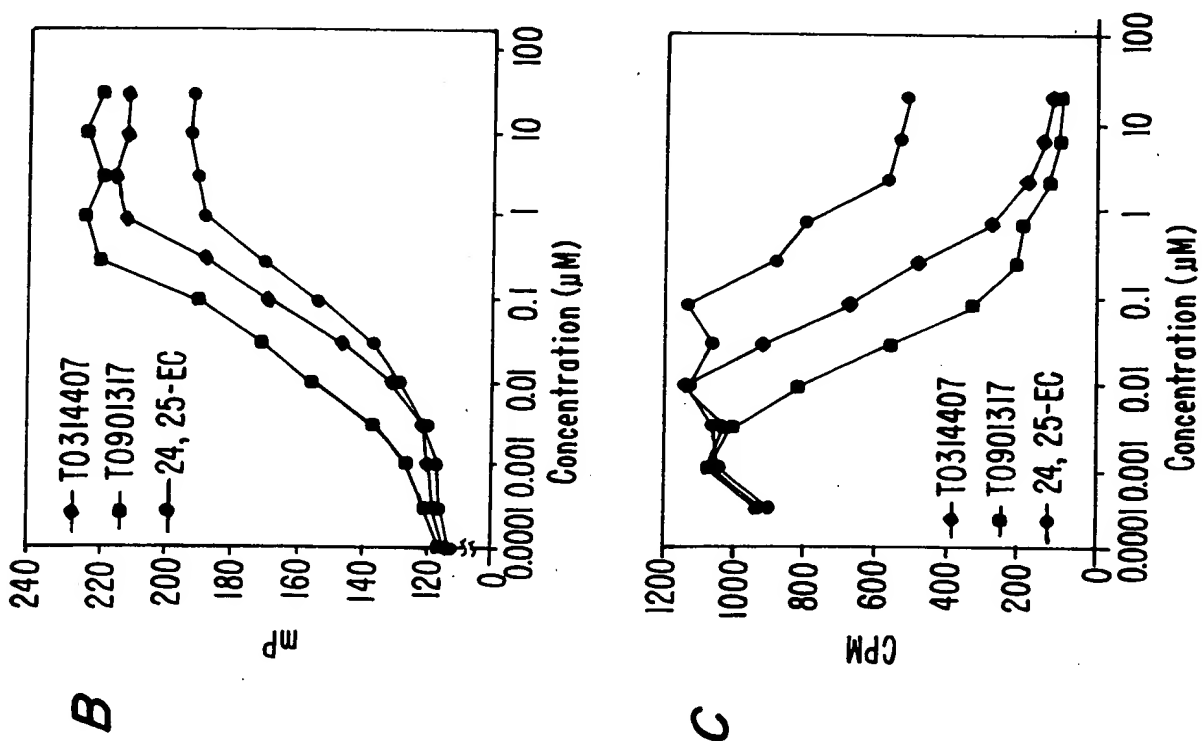


FIG. 6.



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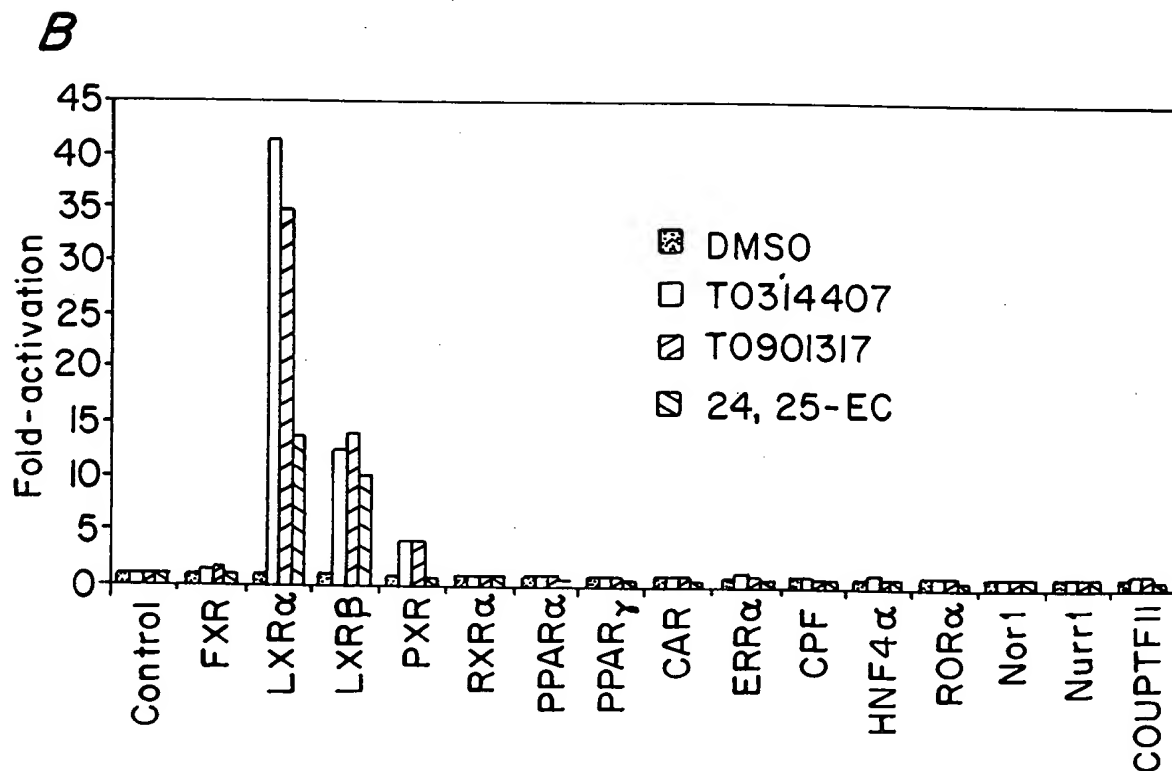
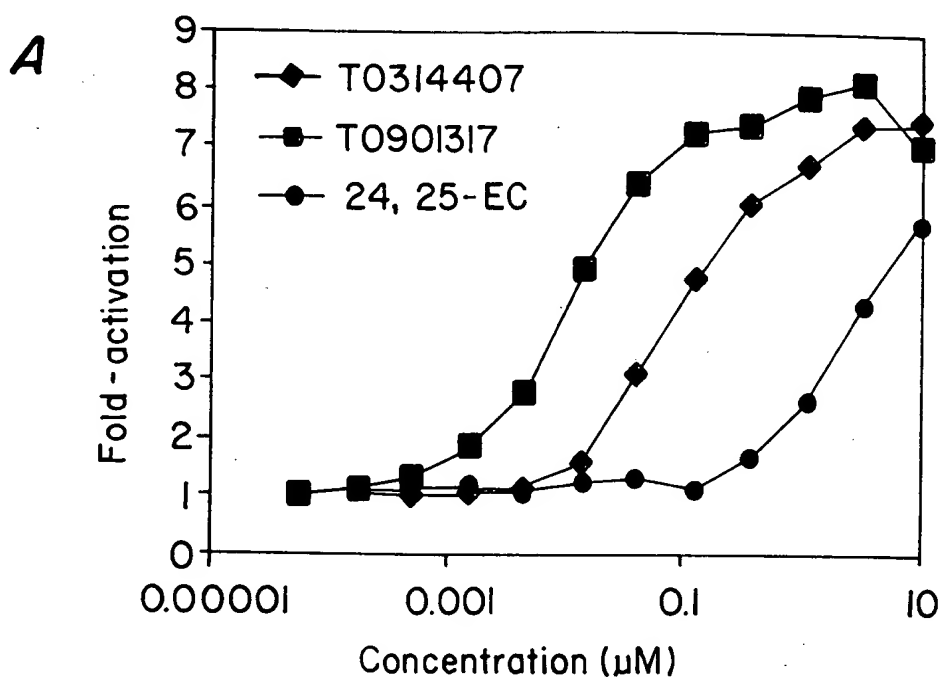


FIG. 7.

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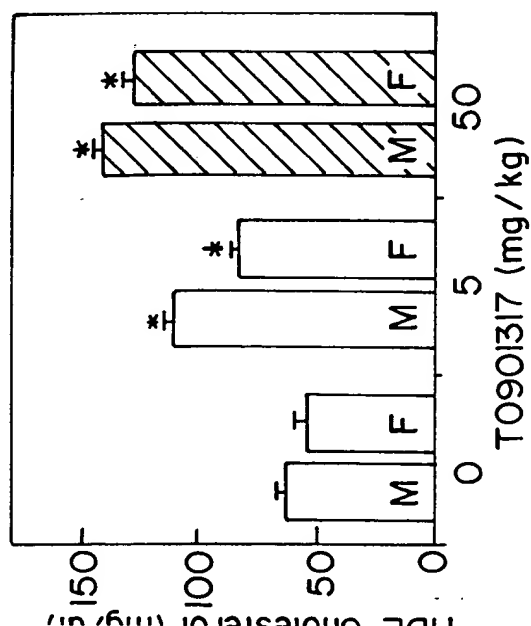
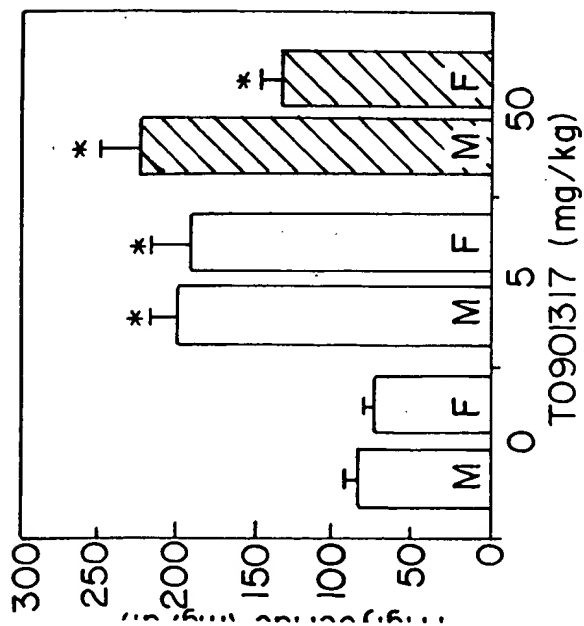
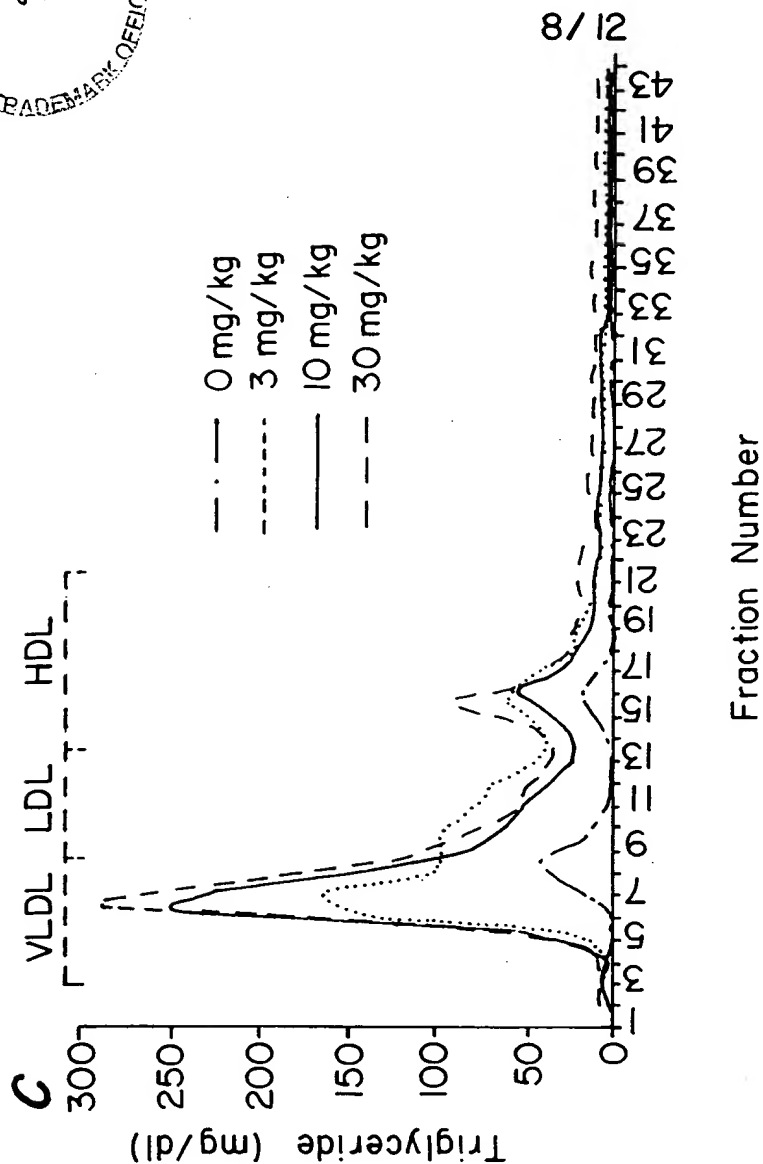


FIG. 8.

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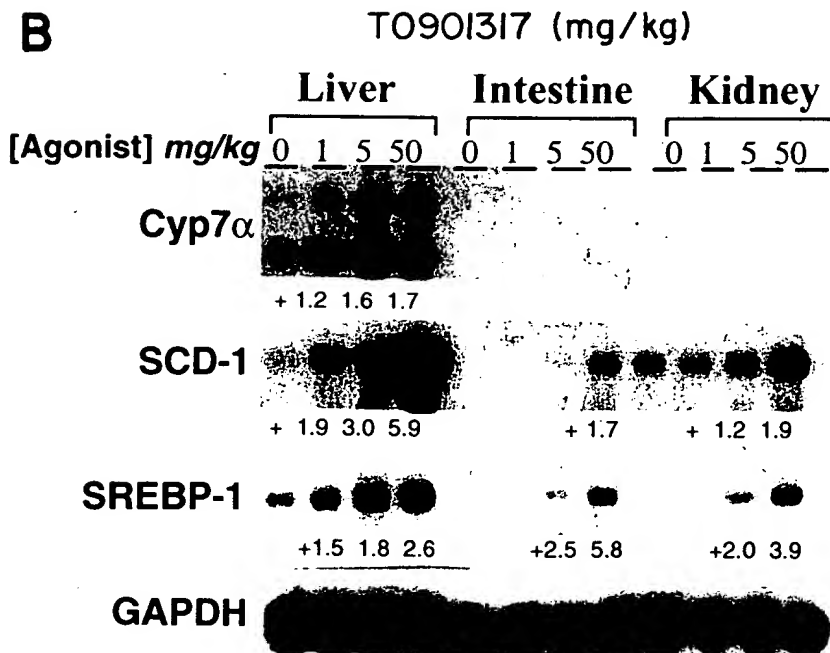
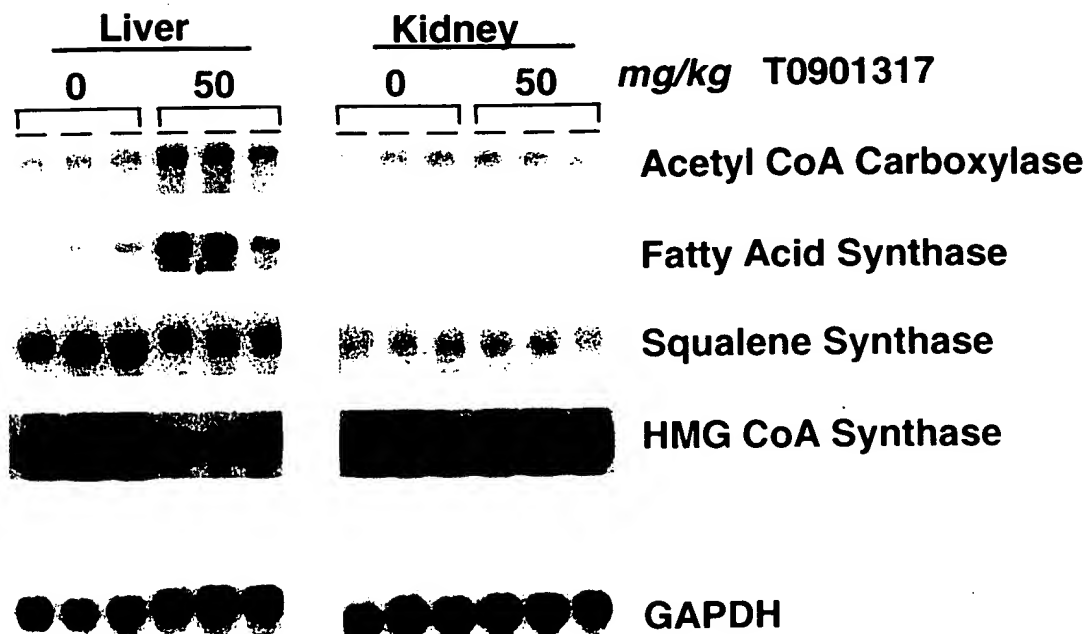


FIG. 9

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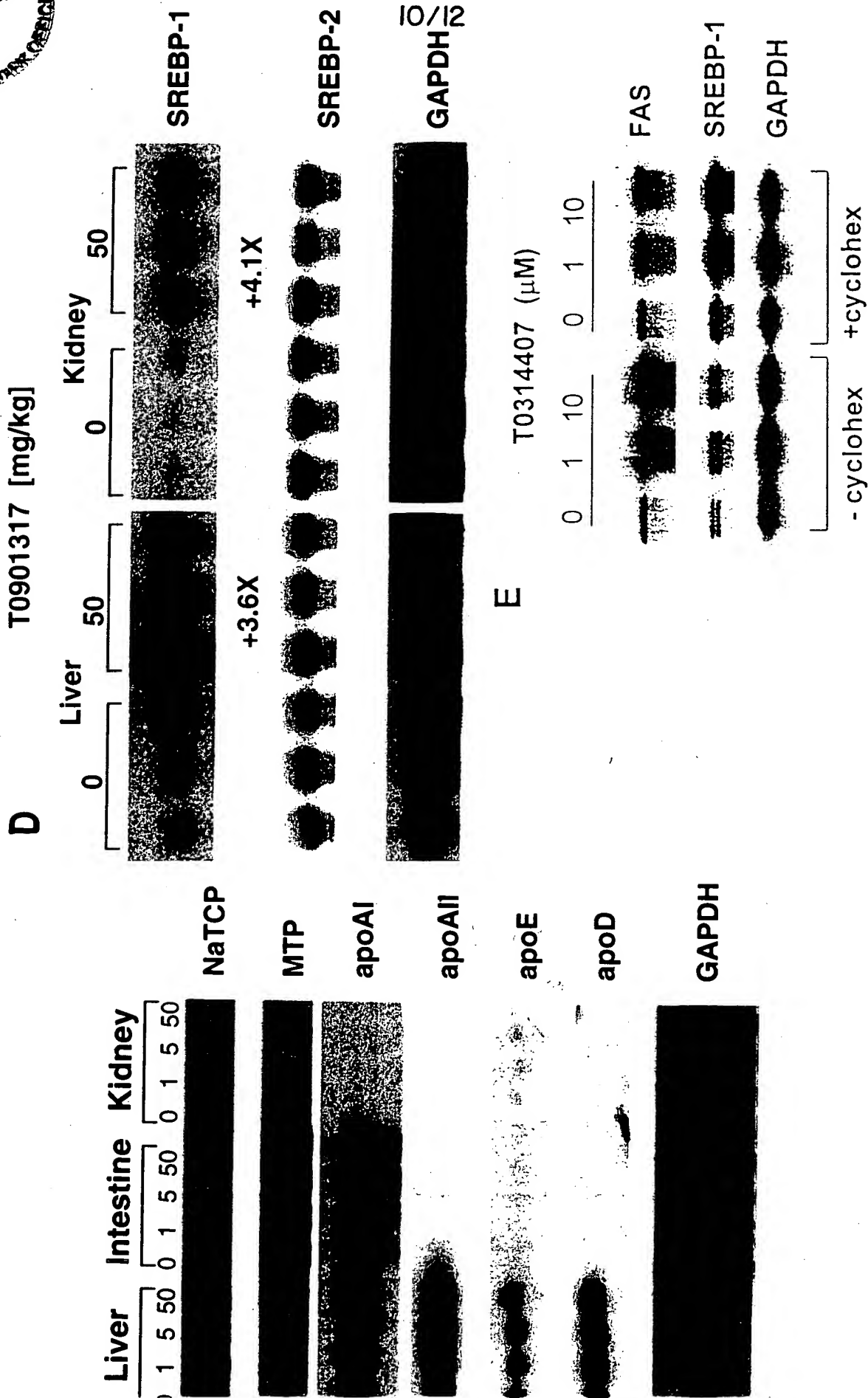


FIG. 9. (Continued)

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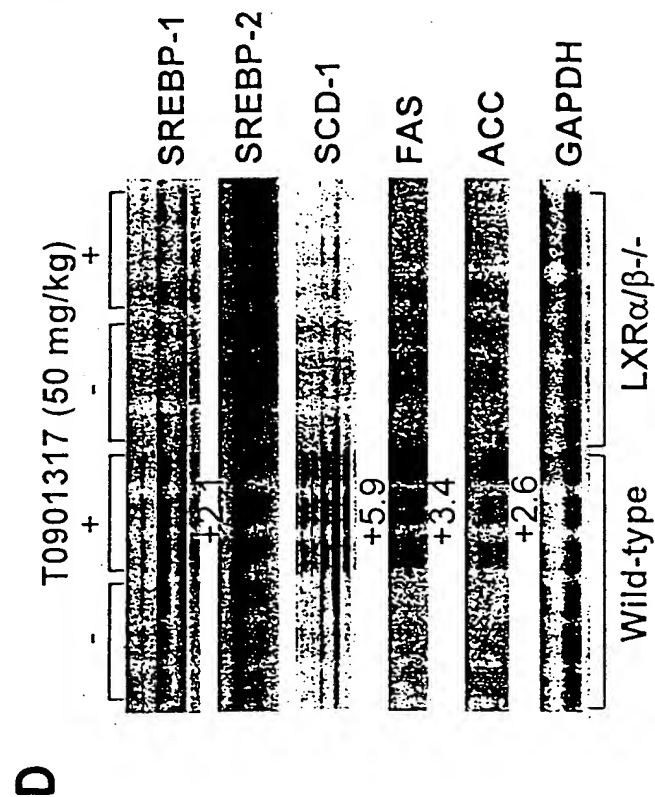
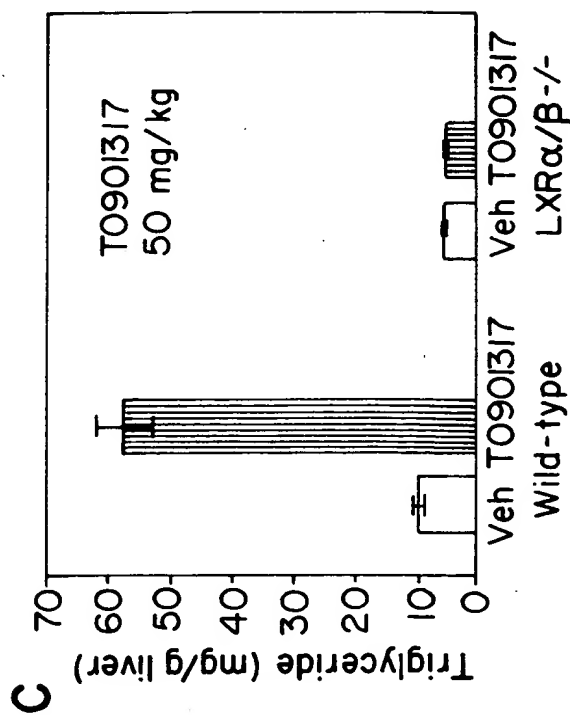
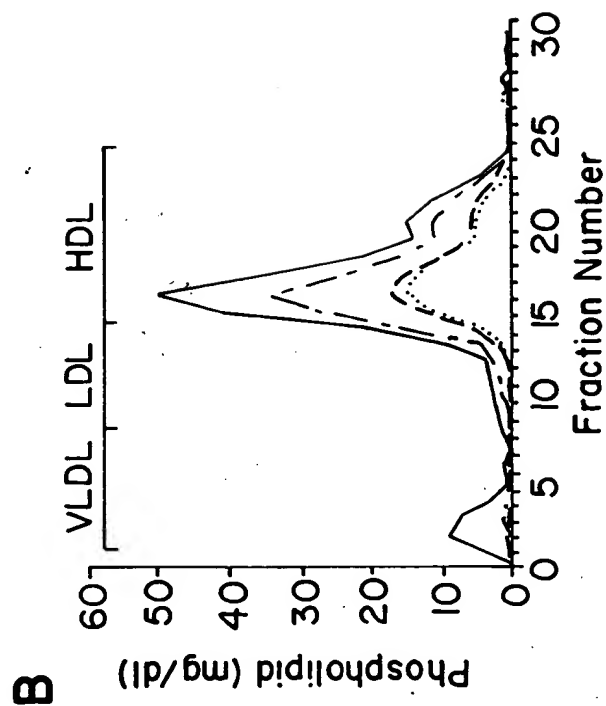
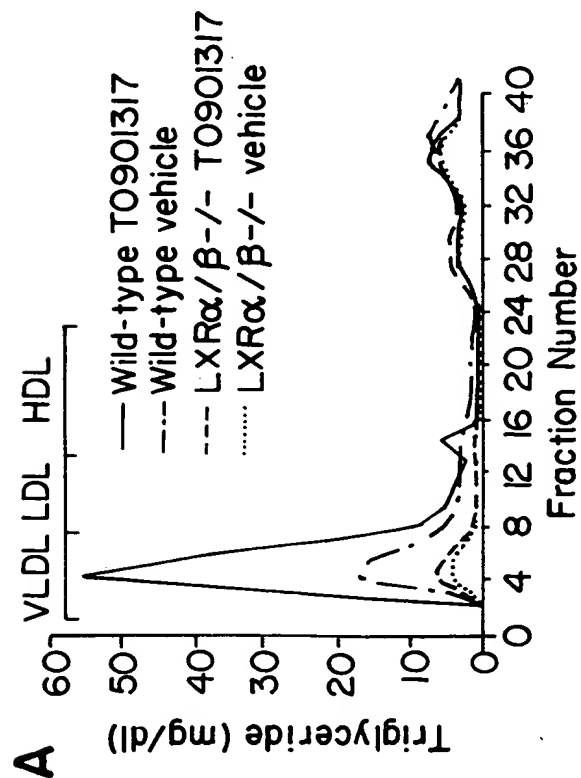
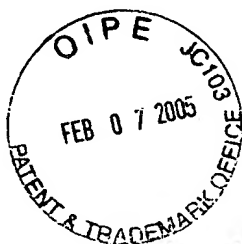


FIG. 10.



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3DNA Probe	Primer Set	Primer Sequence	PCR Product Length
etyl CoA carboxylase	5'	TACCTGTGGGACAGCAACCA	521
	3'	TGGTCGACAGCAAGGGAA	
ty acid synthase	5'	CCCGATGGGGTTTCATC	493
	3'	GATCCCTCTGGGTGTCCC	
ualene synthase	5'	CGTGCAGTGCTTGAATGAACT	288
	3'	TTGGGGATCCGGTGATAAAT	
MG CoA synthase	5'	CCCAGCAGAGGTTTCTACAA	1285
	3'	AATTCCCTCAGGGGACATGC	
REBP-1	5'	TCA ACAACCAAGACAGTGACTTCCCTGGCC	1028
	3'	GTTCTCCTGCTTGAGCTTCTGGTTGCTGTG	
CP	5'	AGCAAGATCAAGGCTCACTTCT	405
	3'	ATAGTGTGGCCTTTGGACTTC	
TP	5'	TTCTCTGCTTCTTCTCCTCCTA	403
	3'	GGCTCGTTTTTCATAGGAGTAGA	
oA-I	5'	GGCAGAGACTATGTGTCCAGTTTGA	564
	3'	GTCATCCAGCGGGTTTGGCCTTCTC	
oA-II	5'	ATAGTCTGCCATCATGAAGCTG	406
	3'	GAGAAACAGGCAGAGGTTAGG	
oE	5'	CCGTGCTGTTGGTCACATT	1014
	3'	TTATTAAGCAAGGGCCACCA	

FIG. 11.